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## Exhibit B

☐ 1: NP\_195636. Reports ...[gi:15234999] The record has been replaced by 145356202

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LOCUS NP\_195636 97 aa linear PLN 09-JUN-2006  
DEFINITION DNA binding / transcription factor [Arabidopsis thaliana].  
ACCESSION NP\_195636  
VERSION NP\_195636.1 GI:15234999  
DBSOURCE REFSEQ: accession NM\_120086.1  
KEYWORDS .  
SOURCE Arabidopsis thaliana (thale cress)  
ORGANISM Arabidopsis thaliana  
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;  
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.  
COMMENT PROVISIONAL REFSEQ: This record has not yet been subject to final  
NCBI review. The reference sequence was derived from AT4G39250.1p.  
[WARNING] On Apr 18, 2007 this sequence was replaced by  
gi:145356202.  
Method: conceptual translation.  
FEATURES  
source 1..97  
/organism="Arabidopsis thaliana"  
/db\_xref="taxon:3702"  
/chromosome="4"  
/ecotype="Columbia"  
Protein 1..97  
/product="DNA binding / transcription factor"  
/calculated\_mol\_wt=10697  
Region 14..60  
/region\_name="SANT"  
/note="'SWI3, ADA2, N-CoR and TFIIIB' DNA-binding domains.  
Tandem copies of the domain bind telomeric DNA tandem  
repeats as part of the capping complex. Binding is sequence  
dependent for repeats which contain the G/C rich motif  
[C2-3 A (CA)1-6]; cd00167"  
Site order(14,47..48,50..51,53..55,57..59)  
/site\_type="DNA binding"  
/db\_xref="CDD:29107"  
CDS 1..97  
/locus\_tag="AT4G39250"  
/coded\_by="NM\_120086.1:1..294"  
/GO\_component="chloroplast; nucleus"  
/GO\_function="DNA binding; transcription factor activity  
[PMID 11118137]"  
/GO\_process="regulation of transcription, DNA-dependent"  
/note="myb family transcription factor, contains PFAM  
profile: PF00249 myb-like DNA binding domain"  
/db\_xref="GeneID:830081"  
/db\_xref="TAIR:AT4G39250"  
ORIGIN  
1 masssmssqs sgswtakqnk afeqalatyd qdtpnrwqnv akvvggktte evkrhyellv

61 qdinsiengh vpfpnyrtsg gctngrlsqe ekryvls  
//

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Aug 28 2007 16:53:42